## RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 69

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/890,684B

DATE: 04/11/2005 TIME: 13:57:22

Input Set : N:\FATIMA\09890684.txt

Output Set: N:\CRF4\04112005\1890684B.raw

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3 <110> APPLICANT: Bullerdiek, Jorn
     5 <120> TITLE OF INVENTION: Preparation for the Prevention and/or Treatment of a Tissue
Change of
      6
             Mesenchymal Origin
     7 <130> FILE REFERENCE: BOH6278P0010US
     9 <140> CURRENT APPLICATION NUMBER: 09/890,684B
     10 <141> CURRENT FILING DATE: 2001-08-03
    12 <150> PRIOR APPLICATION NUMBER: PCT/DE00/00364
    13 <151> PRIOR FILING DATE: 2000-02-04
    15 <160> NUMBER OF SEQ ID NOS: 47
    17 <170> SOFTWARE: PatentIn version 3.1
    19 <210> SEQ ID NO: 1
    20 <211> LENGTH: 19
    21 <212> TYPE: DNA
    22 <213> ORGANISM: Artificial
    24 <220> FEATURE:
    25 <221> NAME/KEY: misc_feature
    26 <223> OTHER INFORMATION: primer HsgA1
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34 <211> LENGTH: 14

35 <212> TYPE: DNA

36 <213 > ORGANISM: Artificial

38 <220> FEATURE:

39 <221> NAME/KEY: misc\_feature

40 <223> OTHER INFORMATION: primer HsgA2

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44 acggttactt kttt 47 <210> SEQ ID NO: 3

48 <211> LENGTH: 18

49 <212> TYPE: DNA

50 <213> ORGANISM: Artificial

52 <220> FEATURE:

53 <221> NAME/KEY: misc\_feature

54 <223> OTHER INFORMATION: primer HagBl

57 <400> SEQUENCE: 3

58 tctattccct acctggat

61 <210> SEQ ID NO: 4 62 <211> LENGTH: 17

63 <212> TYPE: DNA

64 <213 > ORGANISM; Artificial

66 <220> FEATURE:

67 <221> NAME/KEY misc\_feature

14

18

RAW SEQUENCE LISTING DATE: 04/11/2005 PATENT APPLICATION: US/09/890,684B TIME: 13:57:22

Input Set : N:\FATIMA\09890684.txt

Output Set: N:\CRF4\04112005\1890684B.raw

| 68  | <223> OTHER INFORMATION: primer HsgB2                             |    |
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| 71  | <400> SEQUENCE: 4   |    |
| 72  | actettaacg geagtag  | 17 |
|     | <210> SEQ ID NO: 5  |    |
|     | <211> LENGTH: 32  |    |
|     | <212> TYPE: DNA   |    |
| 78  | <213> ORGANISM: Artificial  |    |
|     | <220> FEATURE:  |    |
| 81  | <221> NAME/KEY: misc_feature                                      |    |
| 82  | <223> OTHER INFORMATION: ADE1Bg12S                                |    |
|     | <400> SEQUENCE: 5   |    |
| 86  | gaagatettt atagatggaa tggtgecaac at                               | 32 |
|     | <210> SEQ ID NO: 6  |    |
|     | <211> LENGTH: 31  |    |
|     | <212> TYPE: DNA   |    |
|     | <213> ORGANISM: Artificial  |    |
|     | <220> FEATURE:  |    |
| 95  | <221> NAME/KEY: misc_feature                                      |    |
| 96  | <223> OTHER INFORMATION: primer ADE1Hi3AS                         |    |
|     | <400> SEQUENCE: 6   |    |
| 100 | cccaagctta aaactcttct cgctggcagt c                                | 31 |
|     | <210> SEQ ID NO: 7  |    |
|     | <211> LENGTH: 17<br><212> TYPE: DNA                               |    |
|     | <pre>&lt;212&gt; IIFE: DNA &lt;213&gt; ORGANISM: Artificial</pre> |    |
|     | <220> FEATURE:  |    |
|     | <pre>&lt;221&gt; NAME/KEY: misc_feature</pre>                     |    |
| 110 | <223> OTHER INFORMATION: primer HsqC1                             |    |
| 113 | <pre>&lt;400&gt; SEQUENCE: 7</pre>                                |    |
| 114 | acctttgact cttctgt  |    |
|     | <210> SEQ ID NO: 8  | 17 |
|     | <211> LENGTH: 17  |    |
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|     | <221> NAME/KEY: misc_feature                                      |    |
| 124 | <223> OTHER INFORMATION: primer HsgC2                             |    |
| 127 | <400> SEQUENCE: 8   |    |
|     | tecttgtatt tagtate  |    |
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| 132 | <211> LENGTH: 18  |    |
| 133 | <212> TYPE: DNA   |    |
|     | <213> ORGANISM: Artificial  |    |
| 136 | <220> FEATURE:  |    |
|     | <221> NAME/KEY: misc_feature                                      |    |
| 138 | <223> OTHER INFORMATION: primer HagDl                             |    |
| 141 | <400> SEQUENCE: 9   |    |
| 142 | ccatcatgtt cgactcct   | •  |
| 145 | <210> SEQ ID NO: 10   | 18 |
|     | • •   |    |
|     |   |    |

RAW SEQUENCE LISTING DATE: 04/11/2005
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Input Set : N:\FATIMA\09890684.txt

Output Set: N:\CRP4\04112005\1890684B.raw

146 <211> LENGTH: 17 147 <212> TYPE: DNA 148 <213> ORGANISM: Artificial 150 <220> FEATURE: 151 <221> NAME/KEY: misc\_feature 152 <223> OTHER INFORMATION: primer HagD2 155 <400> SEQUENCE: 10 156 aggtagccgg tgaagcc 17 159 <210> SEQ ID NO: 11 160 <211> LENGTH: 18 161 <212> TYPE: DNA 162 <213> ORGANISM: Artificial 164 <220> FEATURE: 165 <221> NAME/KEY: misc\_feature 166 <223> OTHER INFORMATION: HsgEl 169 <400> SEQUENCE: 11 170 gactetteeg teagetgg 18 173 <210> SEQ ID NO: 12 174 <211> LENGTH: 17 175 <212> TYPE: DNA 176 <213> ORGANISM: Artificial 178 <220> FEATURE: 179 <221> NAME/KEY: misc feature 180 <223> OTHER INFORMATION: primer HsgE2 183 <400> SEQUENCE: 12 184 gctggtaacg gcgctct 17 187 <210> SEQ ID NO: 13 188 <211> LENGTH: 17 189 <212> TYPE: DNA 190 <213> ORGANISM: Artificial 192 <220> FEATURE: 193 <221> NAME/KEY: misc\_feature 194 <223 > OTHER INFORMATION: primer HsgF1 197 <400> SEQUENCE: 13 198 atttctattc cttcgcg 17 201 <210> SEQ ID NO: 14 202 <211> LENGTH: 17 203 <212> TYPE: DNA 204 <213> ORGANISM: Artificial 206 <220> FEATURE: 207 <221> NAME/KEY: misc\_feature 208 <223> OTHER INFORMATION: primer HsgF2 211 <400> SEQUENCE: 14 212 teaggettgg taeggee 17 215 <210> SEQ ID NO: 15 216 <211> LENGTH: 430 217 <212> TYPE: DNA 218 <213 > ORGANISM: Adenovirus

220 <220> FEATURE:

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/890,684B

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TIME: 13:57:22
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Input Set : N:\FATIMA\09890684.txt

Output Set: N:\CRF4\04112005\1890684B.raw

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221 <221> NAME/KEY: misc_feature
 222 <223> OTHER INFORMATION: isolate X765551Ko
 225 <400> SEQUENCE: 15
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 228 agetggeetg geaatgacag getgttgage ccaaatgagt ttgaaatcaa gegeaetgtg
                                                                            60
                                                                           120
 230 gacggggaag ggtacaatgt ggcccaatgt aacatgacca aagactggtt cctggttcag
 232 atgettgeca actacaacat tggctaccag ggettttaca teectgaggg atacaaggat
                                                                           180
 234 cgcatgtact cctttttcag aaacttccag cctatgagca ggcaggtggt tgatgaggtt
                                                                           240
                                                                           300
 236 aattacactg actacaaagc cgtcacctta ccataccaac acaacaactc tggctttgta
                                                                           360
 238 gggtatettg cacetactat gagacaaggg gaacettace cagecaatta tecataceeg
                                                                           420
 240 ctcatcggaa
                                                                           430
 243 <210> SEQ ID NO: 16
 244 <211> LENGTH: 430
 245 <212> TYPE: DNA
246 <213> ORGANISM: Adenovirus
248 <220> FEATURE:
249 <221> NAME/KEY: misc_feature
250 <223> OTHER INFORMATION: isolate M2-3s
253 <400> SEQUENCE: 16
254 ggcacctttt accttaacca cactttcaag aaggtctcca tcatgtttga ctcctcagtc
256 agetggeetg geaatgaeag getgttgage ceaaatgagt ttgaaateaa gegeaetgtg
                                                                            60
258 gacggggaag ggtacaatgt ggcccaatgt aacatgacca aagactggtt cctggttcag
                                                                           120
260 atgettgeca actacaacat tggetaccag ggettttaca teeetgaggg atacaaggat
                                                                           180
262 cgcatgtact cetttttcag aaacttccag cetatgagca ggcaggtggt tgatgaggtt
                                                                           240
264 aattacactg actacaaagc cgtcacctta ccataccaac acaacaactc tggctttgta
                                                                           300
266 gggtatettg cacctactat gagacaaggg gaaccttacc cagccaatta tccatacccg
                                                                          360
                                                                          420
268 ctcatcggaa
                                                                           430
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272 <211> LENGTH: 430
273 <212> TYPE: DNA
274 <213> ORGANISM: Adenovirus
276 <220> FEATURE:
277 <221> NAME/KEY: misc_feature
278 <223> OTHER INFORMATION: isolate M7-1s
281 <400> SEQUENCE: 17
282 ggcaccttct accttaacca cactttcaag aaggtctcca tcatgtttga ctcctcagtc
284 agetggeetg geaatgacag getgttgage ceaaatgagt ttgaaatcaa gegeaetgtg
                                                                           60
                                                                          120
286 gacggggaag gatacaacgt ggcacaatgc aacatgacca aagactggtt cctagttcag
288 atgettgeea actacaacat tggetaccag ggettitaca teeetgaggg atacaaggat
                                                                          180
                                                                          240
290 cgcatgtact ctttttcag aaacttccag cctatgagca ggcaggtggt tgatgaggtt
                                                                          300
292 aattacactg actacaaagc cgtcacctta ccataccaac acaacaacte tggctttgta
                                                                          360
294 gggtaccttg cacctactat gagacaaggg gaaccttacc cagccaatta tccatacccg
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296 ctcatcggaa
                                                                          430
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301 <212> TYPE: DNA
302 <213> ORGANISM: Adenovirus
304 <220> FEATURE:
305 <221> NAME/KEY: misc_feature
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60

120

180

240 300

360

420

430

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RAW SEQUENCE LISTING
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Input Set : N:\FATIMA\09890684.txt
Output Set: N:\CRF4\04112005\1890684B.raw

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306 <223> OTHER INFORMATION: isolate M8-2s
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 310 ggcaccttct accttaacca cactttcaag aaggtctcca tcatgtttga ctcctcagtc
 312 agetggeetg geaatgaeag getgttgage ceaaatgagt ttgaaateaa gegeaetgtg
 314 gacggggaag gatacaacgt ggcacaatgc aacatgacca aagactggtt cctagttcag
 316 atgettgeca actacaacat tggetaccag ggettttaca teeetgaggg atacaaggat
 318 cgcatgtact ctttttcag aaacttccag cctacgagca ggcaggtggt tgatgaggtt
 320 aattacactg actacaaagc cgtcacctta ccataccaac acaacaactc tggctttgta
 322 gggtacettg cacetactat gagacaaggg gaacettace eggecaatta tecataceeg
 324 ctcatcggaa
 327 <210> SEQ ID NO: 19
 328 <211> LENGTH: 143
 329 <212> TYPE: PRT
 330 <213> ORGANISM: Adenovirus: isolate X765551Ko
 332 <400> SEQUENCE: 19
333 Gly Thr Phe Tyr Leu Asn His Thr Phe Lys Lys Val Ser Ile Met Phe
336 Asp Ser Ser Val Ser Trp Pro Gly Asn Asp Arg Leu Leu Ser Pro Asn
337
                 20
339 Glu Phe Glu Ile Lys Arg Thr Val Asp Gly Glu Gly Tyr Asn Val Ala
342 Gln Cys Asn Met Thr Lys Asp Trp Phe Leu Val Gln Met Leu Ala Asn
        5 Q
                              55
345 Tyr Asn Ile Gly Tyr Gln Gly Phe Tyr Ile Pro Glu Gly Tyr Lys Asp
                         70
                                              75
348 Arg Met Tyr Ser Phe Phe Arg Asn Phe Gln Pro Met Ser Arg Gln Val
                     85
                                          90 .
351 Val Asp Glu Val Asn Tyr Thr Asp Tyr Lys Ala Val Thr Leu Pro Tyr
                                     105
354 Lys His Asn Asn Ser Gly Phe Val Gly Tyr Leu Ala Pro Thr Met Arg
355
            115
                                120
357 Gln Gly Glu Pro Tyr Pro Ala Asn Tyr Pro Tyr Pro Leu Ile Gly
358
        130
                            135
361 <210> SEQ ID NO: 20
362 <211> LENGTH: 143
363 <212> TYPE: PRT
364 <213> ORGANISM: Adenovirus: isolate M2-3s
366 <400> SEQUENCE: 20
367 Gly Thr Phe Tyr Leu Asn His Thr Phe Lys Lys Val Ser Ile Met Phe
                      5
                                         10
370 Asp Ser Ser Val Ser Trp Pro Gly Asn Asp Arg Leu Leu Ser Pro Asn
                 20
                                     25
373 Glu Phe Glu Ile Lys Arg Thr Val Asp Gly Glu Gly Tyr Asn Val Ala
             35
                                 40
376 Gln Cys Asn Met Thr Lys Asp Trp Phe Leu Val Gln Met Leu Ala Asn
        50
                             55
379 Tyr Abn Ile Gly Tyr Gln Gly Phe Tyr Ile Pro Glu Gly Tyr Lys Asp
380 65
                         70
382 Arg Met Tyr Ser Phe Phe Arg Asn Phe Gln Pro Met Ser Arg Gln Val
```

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/11/2005 PATENT APPLICATION: US/09/890,684B TIME: 13:57:23

Input Set : N:\FATIMA\09890684.txt

Output Set: N:\CRF4\04112005\1890684B.raw

#### Please Note:

Use of n and/or Kaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Kaa.

Seq#:28; N Pos. 147,148,222,299 Seq#:32; Xaa Pos. 49,50,74,100

#### Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/890,684B

DATE: 04/11/2005 TIME: 13:57:23

Input Set : N:\PATIMA\09890684.txt

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L:7 M:283 W: Missing Blank Line separator, <130> field identifier L:654 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:120

M:341 Repeated in SeqNo=28

L:797 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:48

M:341 Repeated in SeqNo=32

**衛001** 

RECEPTION OK

TX/RX NO

RECIPIENT ADDRESS

DESTINATION ID

ST. TIME TIME USE

PGS. RESULT 8290

703 308 4221

04/11 14:39

02'34

9

OK





### United States Patent and Trademark Scientific and Technical Information Center Biotechnology Systems Branch

1 1 APR 2005

Legal Staff International Division

# FAX TRANSMISSION COVER SHEET

| DATE: _  | 4-11-05     | Total Number of Pages Faxed: |  |  |  |
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| TO:  |             |                              |  |  |  |
| NAME:  | MR. RICHARD | eoss                         |  |  |  |
| ORG.:  | PCT LEBAL   |                              |  |  |  |
| FAX NUMBER: 571-273-0459   |             |                              |  |  |  |
|  | ·           |                              |  |  |  |
|  |             |                              |  |  |  |
| FROM: Mark Spencer<br>Voice Ph. Number: (571)272-2510<br>FAX Ph. Number: (571)273-0221 |             |                              |  |  |  |
| Message: 09/890684B  |             |                              |  |  |  |
|  |             |                              |  |  |  |
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